

Thu, Mar 15 11:53:15 2001

us

Db 310 ATGCTGTACTTTAAAGTTGGTAATGAACACCCCTAAGTATATAAATCACAACCTATAACAGAT 251  
QY 481 gtttaagcctacagatgttaggagttcttagatgaacaaagggttaaagataaacaatttaaca 540  
Db 250 GTTAAGCCTACAGATGTAGGAGTTCTAGATGAACAAAGGCTAAAGATAAACCAATTAAACA 191  
QY 541 ttaattacttctgtagattacaaatgaagaacagcggtttgggaaaaacgtaaaatcttt 600  
Db 150 TTAATTACTTGTGATGATTACATATGAAGAAGACAGCGCTTGGGAAAAAACGTAAATCTTT 131  
QY 601 gtagctacagaagtcataataa 621  
Db 130 GTAGCTACAGAAGTCARATAA 110

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Query Match      100.0%; Score 621; DB 11; Length 3733;
Best local similarity 100.0%; Pred. No. 1.2e-106;
Matches 621; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1  atgaaaatgagcaaatcgattaatgacaaatcgctggtgtgctacttcttcaggcgca 60
DB      730  ATGAAAAAATGGACAAATCGAATTAATGACAATCGCTGCTGTGTAATCTTCTAGTGCCA 671

QY      61  gcaattgtgttgcaaacccacatcgcataattcttcacgatacaagatacaagatgaa 120
DB      670  GCATATTGTTGCTAAACCACATATCGATAATTCATCTTCAGATAAAGATAAAGATGAA 611

QY      121  aagattgaacaatcgtacaaagttaaagaacaggcgagtaaagataaaaagcagcaaa 180
DB      610  AAGATTGAACAAATATGATAAAATGTAAAGAACAGCGGAGTAAAGATATAAAGCAGCAA 551

QY      181  gctaaacctcaaatccgaagataaaatcgaagatggcggtcatattgaaattccagat 240
DB      550  GCTAAACCTCNAAATCCGAAAGATAAATCGAAAGTGGCAGGCTATATTGAAATCCAGAT 491

QY      241  gctgatattaagaagaccgttatccaggaccagcaacacotgcaacaattcaatagaggt 300
DB      490  GCTGATATTAAAGAACCCAGTATATCCAGGACCGAGCAACCTGCAACATTAATATAGAGGT 431

QY      301  gtaagctttgcagagaaaaatgaaatcactagatgatacaaaatatttcaattgcaggacac 360
DB      430  GTAAGCTTTGCAGAGAAAATGAATCACTAGATGATCAAAATATTTCATTCGAGGACAC 371

QY      361  acttcaattgacgtccggaactatcaattcacaattcttaagcagcgccaaaaaagtagt 420
DB      370  ACTTTCATTGACGTCGCCAAGCTATCAATTTACAAATCTTAAAGCAGCAGCAAAAAGTAGT 311

QY      421  atggctgactttcaagagttggtaatgaaacacgcgtcaagatacaaatgacgaagtataaagat 480
DB      —

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RESULT 2  
US-08-781-986A-538/c  
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